

*Holland**Li-R(1)*

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/830,703

DATE: 12/12/2001
 TIME: 11:51:43

Input Set : A:\seqlist.txt
 Output Set: N:\CRF3\12112001\I830703.raw

4 <110> APPLICANT: Biofrontera Pharmaceuticals AG
 5 Lubbert, Hermann
 7 <120> TITLE OF INVENTION: TRANSGENIC ANIMAL MODEL FOR
 8 NEURODEGENERATIVE DISEASES
 10 <130> FILE REFERENCE: STERN1.001APC
 12 <140> CURRENT APPLICATION NUMBER: 09/830,703
 C--> 13 <141> CURRENT FILING DATE: 2001-12-12
 15 <150> PRIOR APPLICATION NUMBER: EP 99116766.9
 16 <151> PRIOR FILING DATE: 1999-08-30
 18 <160> NUMBER OF SEQ ID NOS: 34
 20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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 24 <212> TYPE: DNA
 25 <213> ORGANISM: Mus musculus
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 30 cgtgaccat gatagtgttt gtcaggttca actccagcta tggcttcca gtggaggtcg 180
 31 attctgacac cagcatctt cagctcaagg aagtgttgc taagcgacag ggggttccag 240
 32 ctgaccagct gcgtgtgatt tttgccggga aggagttcc gaatcacctg acggttcaaa 300
 33 actgtgacct ggaacaacag agtattgtac acatagtaca gagaccacgg aggagaagtc 360
 34 atgaaacaaa tgcacatcgga ggggacgaac cccagagcac ctcagagggc tccatatggg 420
 35 agtccaggag cttgacacga gtggacactga gcagccatac cctgcccgtg gactctgtgg 480
 36 ggctggcggt cattctggac acagacagta agaggattc agaagcagcc agaggtccag 540
 37 ttaaaccac ctacaacacg ttttcatct actgcaaagg cccctccac aaggtccacg 600
 38 ctggaaagct ccgagttcag tggcacct gcaacaagc aaccctcacc ttggcccagg 660
 39 gcccatctt cttggacat gtcttaattt caaaccggat gatgttgc tgccagtctc 720
 40 cagactgccc tggaaaccaga gctgaatttt tctttaatg tggagcacac ccaacctcag 780
 41 acaaggacac gtcggtagct ttgaacctga tcaccagcaa caggcgcacg atcccttgca 840
 42 tagcgtgcac agatgtcagg agccctgtcc tggcttcca gtgttaaccac cgtcacgtga 900
 43 tctgttttgc ctgtttccac ttgtatttgt tcacaagact caacgatcgg cagttgtcc 960
 44 acgatgctca acttggctac tccctggcggt gtgttagctgg ctgtcccaac tccctgatta 1020
 45 aagagctcca tcacttcagg atccttggag aagagcagta cactaggta cagcagtatg 1080
 46 gggccgagga atgcgtctg caaatggggat gtgtgtgtg ccccccgtct ggctgtggag 1140
 47 ctggactgct acctgaacag gGCCAGAGGA aagtccacgt cgaagggggc aacgcctgg 1200
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ENTERED

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Input Set : A:\seqlist.txt
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61 cctatcttta caaatggtc tgatgagatt acaaccctc tgtgtactaa tcagcttata 2040
62 aaccaagtga gaaccttagga aagctaattt gatggcagac tgcttaaattc gcagggagga 2100
63 ctcagaagcc aaacctactt ccgttcgttt cattatctgc aacttttagaa agaaatgatc 2160
64 ttttttccc cctgaaaaga taacaaagtc tgcaatttgg tttggaggtat tcctactgca 2220
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70 atgcattcaca ctcaagtgtt aatctttcaa gttttcttt tctttttcct gttttttatt 2580
71 tttttgtt ttgctttttt tttttttttt tttgggtgtg gtggggctac caaacttgag 2640
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87 <213> ORGANISM: Mus musculus
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92 tggatgtttt tggatgtttc aactccagct atgggttccc agtggaggtc gattctgaca 180
93 ccagcatctt gcagctcaag gaagtggttt ctaagcgaca gggggttcca gctgaccgc 240
94 tgcgtgttat ttttgcggg aaggagctt cgaatcacct gacggttcaa aactgtgacc 300
95 tggatgtttt ctttgcggg aaggagctt cgaatcacct gacggttcaa aactgtgacc 360
96 atgcattcgg aggggacgaa ccccgagca cctcagaggg ctccatatgg gatgtccagga 420
97 gcttgcacacg agtggacactg agcagccata ccctgcccgt ggactctgtg gggctggcgg 480
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99 ccacccatcaa cagcttttccatctactgca aaggcccctg ccacaagggtc cagctggaa 600
100 agctcccgagt tcagtgtggc acctgcaaaac aagcaaccct caccttgcgc caggcccat 660
101 cttgcgtggaa cgatgtcttta attccaaacc ggtatgtgtt tgatgtccag tctccagact 720
102 gcccctggaaac cagactgtaa tttttcttta aatgtggatc acacccaaacc tcagacaagg 780
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104 gcacagatgt cagtcattt cctctgtcat ctggtgccct cgtgtggact cggcctcatc 900
105 tccactgaac ctgtttctt aggactgtgc aataggttgc caccccttac tgagaacaag 960
106 gcagcttctg gtctcttggc ttccttgcgtt ccaacggcag cattgtactgt acacccttca 1020
107 gtccttaccaa ccccttacc tggatgtgtt ctgttgcgtt tagcttctcc aagatgccta 1080
108 tttccacacaca cagtttcttgcgtt cttttccat ccccccattag gtttatgcgc atgatgttgc 1140
109 accgcacccatc atgatgttgcgtt gtttgcgtata caagacttcc tggatcccc gcttgcgttcc 1200

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110 tagaatcccc tggaaactggg ttcaagtccat tatcttcaat agccctttt aaaagttagt 1260
 111 tcttgggctg gtgagatggc tcagtggta agagcacccg actgcttcc cgaagtccag 1320
 112 agttcaaaaat cccagcaacc acatgggtgc tcacaaccat ccgtaacaag atctgactcc 1380
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 119 <213> ORGANISM: Mus musculus
 121 <400> SEQUENCE: 3
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 124 gttcaactcc agctatggct tcccagtgg agtcgattct gacaccagca tcttgagct 180
 125 caaggaagtg gttgctaagc gacaggggg tccagctgac cagctcggt tgattttgc 240
 126 cgggaaggag cttccgaatc acctgacggt tcaaaaactgt gacctggAAC aacagagtat 300
 127 tgtacacata gtacagagac cacggaggag aagtcatgaa acaaattgtat ctggagggg 360
 128 cgaaccccg agcacccatc agggctccat atggagatcc aggagcttga cagcgtgg 420
 129 cctgagcagc cataccctgc cggtgactc tggggctg gcggtcattt tggacacaga 480
 130 cagtaagagg gattcagaag cagccagagg tccagcgtt aaacccaccc acaacagctt 540
 131 tttcatctac tgcaaaggcc cctgccacaa ggtccagct ggaaagctcc gagttcagt 600
 132 tggcacctgc aaacaagcaa ccctcaccc ggcccaaggc ccatcttgc gggacgatgt 660
 133 cttaaattcca aaccggatga gtggtgagtg ccagtcctca gactgccctg gaaccagagc 720
 134 tgaatttttc tttaaatgtg gagcacaccc aacccatcagac aaggacacgt cggttagctt 780
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 136 tatgcgcattt agttac 857
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 139 <211> LENGTH: 464
 140 <212> TYPE: PRT
 141 <213> ORGANISM: Mus musculus
 143 <400> SEQUENCE: 4
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 146 Val Asp Ser Asp Thr Ser Ile Leu Gln Leu Lys Glu Val Val Ala Lys
 147 20 25 30
 148 Arg Gln Gly Val Pro Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys
 149 35 40 45
 150 Glu Leu Pro Asn His Leu Thr Val Gln Asn Cys Asp Leu Glu Gln Gln
 151 50 55 60
 152 Ser Ile Val His Ile Val Gln Arg Pro Arg Arg Arg Ser His Glu Thr
 153 65 70 75 80
 154 Asn Ala Ser Gly Gly Asp Glu Pro Gln Ser Thr Ser Glu Gly Ser Ile
 155 85 90 95
 156 Trp Glu Ser Arg Ser Leu Thr Arg Val Asp Leu Ser Ser His Thr Leu
 157 100 105 110
 158 Pro Val Asp Ser Val Gly Leu Ala Val Ile Leu Asp Thr Asp Ser Lys
 159 115 120 125
 160 Arg Asp Ser Glu Ala Ala Arg Gly Pro Val Lys Pro Thr Tyr Asn Ser
 161 130 135 140
 162 Phe Phe Ile Tyr Cys Lys Gly Pro Cys His Lys Val Gln Pro Gly Lys

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163 145          150          155          160
164 Leu Arg Val Gln Cys Gly Thr Cys Lys Gln Ala Thr Leu Thr Leu Ala
165          165          170          175
166 Gln Gly Pro Ser Cys Trp Asp Asp Val Leu Ile Pro Asn Arg Met Ser
167          180          185          190
168 Gly Glu Cys Gln Ser Pro Asp Cys Pro Gly Thr Arg Ala Glu Phe Phe
169          195          200          205
170 Phe Lys Cys Gly Ala His Pro Thr Ser Asp Lys Asp Thr Ser Val Ala
171          210          215          220
172 Leu Asn Leu Ile Thr Ser Asn Arg Arg Ser Ile Pro Cys Ile Ala Cys
173 225          230          235          240
174 Thr Asp Val Arg Ser Pro Val Leu Val Phe Gln Cys Asn His Arg His
175          245          250          255
176 Val Ile Cys Leu Asp Cys Phe His Leu Tyr Cys Val Thr Arg Leu Asn
177          260          265          270
178 Asp Arg Gln Phe Val His Asp Ala Gln Leu Gly Tyr Ser Leu Pro Cys
179          275          280          285
180 Val Ala Gly Cys Pro Asn Ser Leu Ile Lys Glu Leu His His Phe Arg
181          290          295          300
182 Ile Leu Gly Glu Glu Gln Tyr Thr Arg Tyr Gln Gln Tyr Gly Ala Glu
183 305          310          315          320
184 Glu Cys Val Leu Gln Met Gly Gly Val Leu Cys Pro Arg Pro Gly Cys
185          325          330          335
186 Gly Ala Gly Leu Leu Pro Glu Gln Gly Gln Arg Lys Val Thr Cys Glu
187          340          345          350
188 Gly Gly Asn Gly Leu Gly Cys Gly Phe Val Phe Cys Arg Asp Cys Lys
189          355          360          365
190 Glu Ala Tyr His Glu Gly Asp Cys Asp Ser Leu Leu Glu Pro Ser Gly
191          370          375          380
192 Ala Thr Ser Gln Ala Tyr Arg Val Asp Lys Arg Ala Ala Glu Gln Ala
193 385          390          395          400
194 Arg Trp Glu Glu Ala Ser Lys Glu Thr Ile Lys Lys Thr Thr Lys Pro
195          405          410          415
196 Cys Pro Arg Cys Asn Val Pro Ile Glu Lys Asn Gly Gly Cys Met His
197          420          425          430
198 Met Lys Cys Pro Gln Pro Gln Cys Lys Leu Glu Trp Cys Trp Asn Cys
199          435          440          445
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205 <211> LENGTH: 262
206 <212> TYPE: PRT
207 <213> ORGANISM: Mus musculus
209 <400> SEQUENCE: 5
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213          20          25          30
214 Arg Gln Gly Val Pro Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys

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215      35          40          45
216 Glu Leu Pro Asn His Leu Thr Val Gln Asn Cys Asp Leu Glu Gln Gln
217      50          55          60
218 Ser Ile Val His Ile Val Gln Arg Pro Arg Arg Arg Ser His Glu Thr
219 65      70          75          80
220 Asn Ala Ser Gly Gly Asp Glu Pro Gln Ser Thr Ser Glu Gly Ser Ile
221      85          90          95
222 Trp Glu Ser Arg Ser Leu Thr Arg Val Asp Leu Ser Ser His Thr Leu
223      100         105         110
224 Pro Val Asp Ser Val Gly Leu Ala Val Ile Leu Asp Thr Asp Ser Lys
225      115         120         125
226 Arg Asp Ser Glu Ala Ala Arg Gly Pro Ala Val Lys Pro Thr Tyr Asn
227      130         135         140
228 Ser Phe Phe Ile Tyr Cys Lys Gly Pro Cys His Lys Val Gln Pro Gly
229 145      150         155         160
230 Lys Leu Arg Val Gln Cys Gly Thr Cys Lys Gln Ala Thr Leu Thr Leu
231      165         170         175
232 Ala Gln Gly Pro Ser Cys Trp Asp Asp Val Leu Ile Pro Asn Arg Met
233      180         185         190
234 Ser Gly Glu Cys Gln Ser Pro Asp Cys Pro Gly Thr Arg Ala Glu Phe
235      195         200         205
236 Phe Phe Lys Cys Gly Ala His Pro Thr Ser Asp Lys Asp Thr Ser Val
237      210         215         220
238 Ala Leu Asn Leu Ile Thr Ser Asn Arg Arg Ser Ile Pro Cys Ile Ala
239 225      230         235         240
240 Cys Thr Asp Val Ser His Leu Pro Leu Ser Ser Gly Ala Ser Val Trp
241      245         250         255
242 Thr Arg Pro His Leu His
243      260
246 <210> SEQ ID NO: 6
247 <211> LENGTH: 250
248 <212> TYPE: PRT
249 <213> ORGANISM: Mus musculus
250 <400> SEQUENCE: 6
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255      20         25          30
256 Arg Gln Gly Val Pro Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys
257      35         40          45
258 Glu Leu Pro Asn His Leu Thr Val Gln Asn Cys Asp Leu Glu Gln Gln
259      50         55          60
260 Ser Ile Val His Ile Val Gln Arg Pro Arg Arg Arg Ser His Glu Thr
261 65      70          75          80
262 Asn Ala Ser Gly Gly Asp Glu Pro Gln Ser Thr Ser Glu Gly Ser Ile
263      85         90          95
264 Trp Glu Ser Arg Ser Leu Thr Arg Val Asp Leu Ser Ser His Thr Leu
265      100        105         110
266 Pro Val Asp Ser Val Gly Leu Ala Val Ile Leu Asp Thr Asp Ser Lys

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/830,703

DATE: 12/12/2001

TIME: 11:51:44

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\12112001\I830703.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

STATISTICS SUMMARY
PATENT APPLICATION: US/09/830,703

DATE: 12/12/2001
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Input Set : A:\seqlist.txt
Output Set: N:\CRF3\12112001\I830703.raw

Application Serial Number: US/09/830,703

Alpha or Numeric: Numeric

Application Class:

Application File Date: 12-12-2001

Art Unit: OIPE

Software Application: FastSeq

Total Number of Sequences: 34

Total Nucleotides: 48922

Total Amino Acids: 3874

Number of Errors: 0

Number of Warnings: 0

Number of Corrections: 1

MESSAGE SUMMARY

271 C: 1 (Current Filing Date differs)